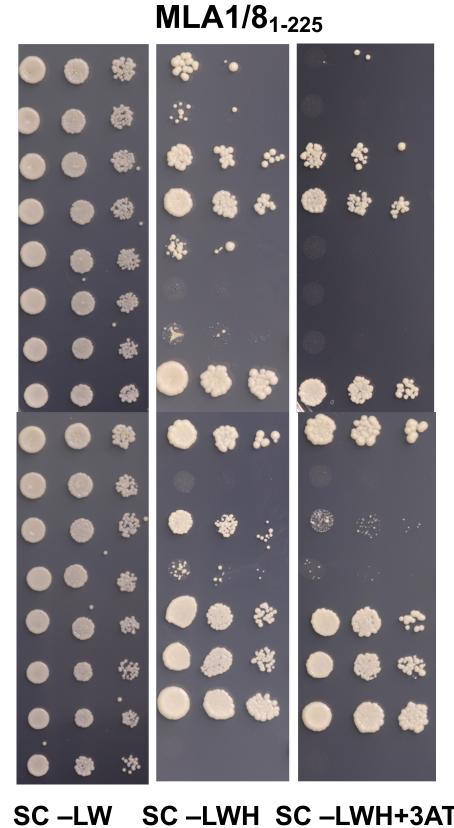
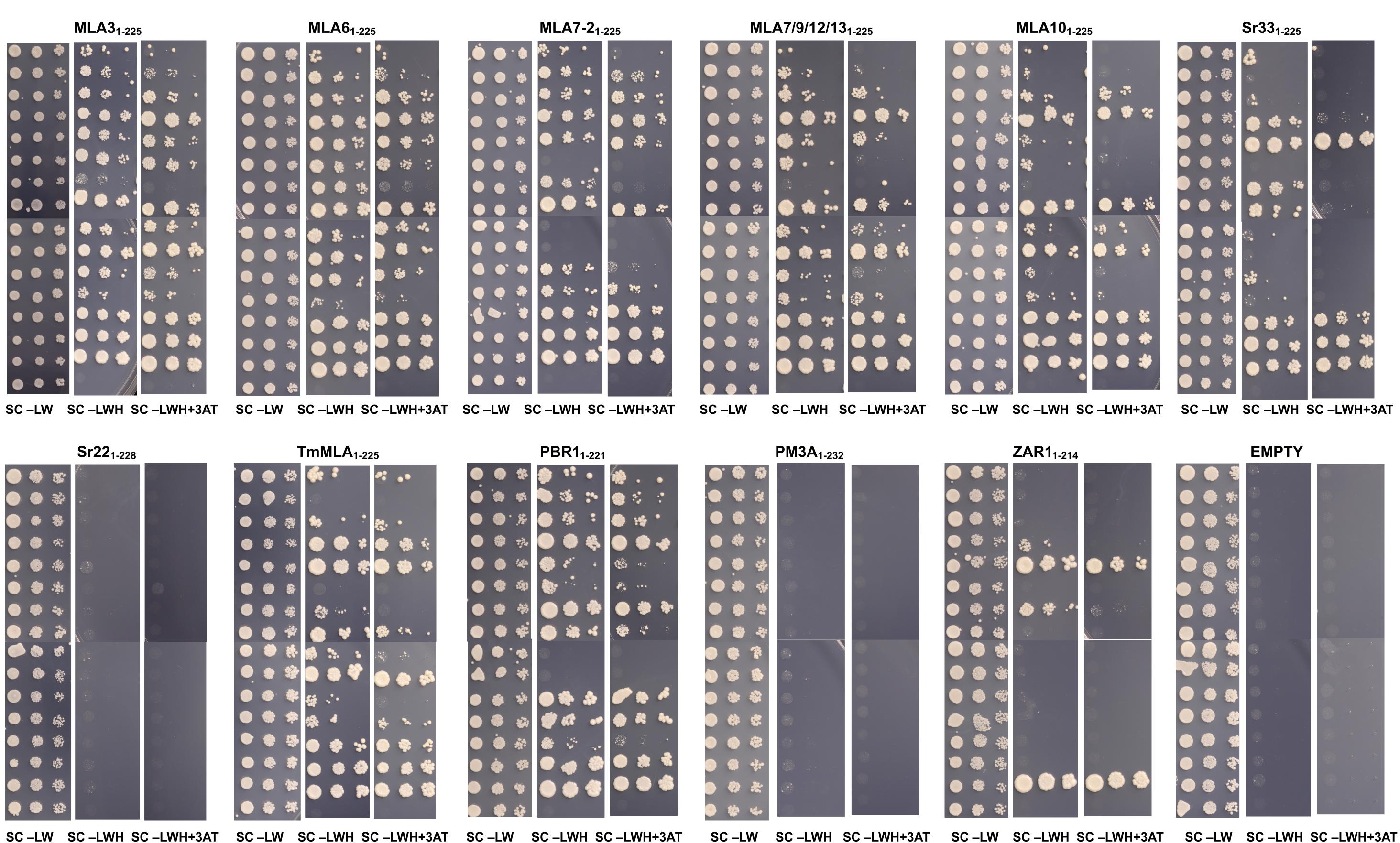
AD Ubiquitn Cue Golgin 5 Dynamin **Kinetochore NDC80** Kinesin PI 4-kinase alpha **OBERON 2-like** Homeobox protein Disulfide isomerase PPR 336 bHLH **POLAR** protein AAA-ATPase 1 WPP protein Polyamine 1-binding **AD-EMPTY** 

BD





Sr50<sub>1-228</sub> Ubiquitn Cue Golgin 5 Dynamin 🕘 🌒 🐟 **Kinetochore NDC80** 🔹 📑 🔒 Kinesin 🕘 🌒 🤹 - 🤹 🔹 PI 4-kinase alpha **OBERON 2-like** s 😽 📩 Homeobox protein . ی ک - - 1 Disulfide isomerase 🕘 🍈 🦂 PPR 336 o 🌸 🏘 bHLH POLAR protein AAA-ATPase 1 WPP protein Polyamine 1-binding AD-EMPTY

SC – LW SC – LWH SC – LWH+3AT SC – LW SC – LWH SC – LWH+3AT

S3 Figure. Y2H validation of interactions for the CC+NB domains of MLA1, MLA3, MLA7-2, MLA8, MLA9, MLA10, MLA12, MLA13; orthologs TmMLA, Sr33 and Sr50; and the outgroup of Sr22, PBR1, PM3A and ZAR1 were cloned into bait vectors and tested for interactions with 15 novel prey candidates. SC-LW media was used as control for diploid growth, and the interaction was tested using selection with SC-LWH and SC-LWH + 0.5 mM 3-AT media.

Velásquez-Zapata et. al. (2022). An interolog-based barley interactome as an integration framework for NLR immune signaling.

